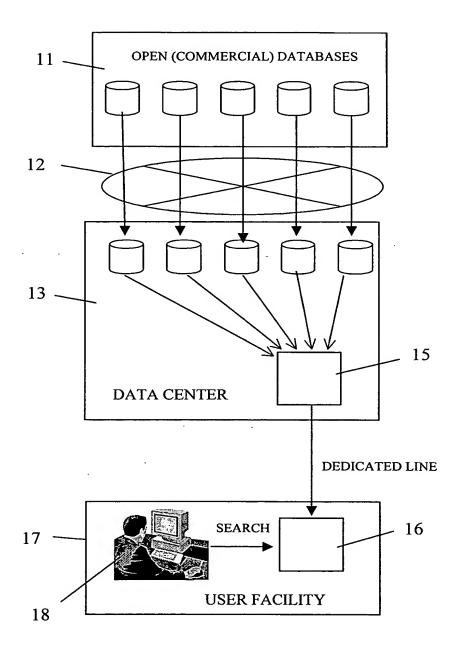
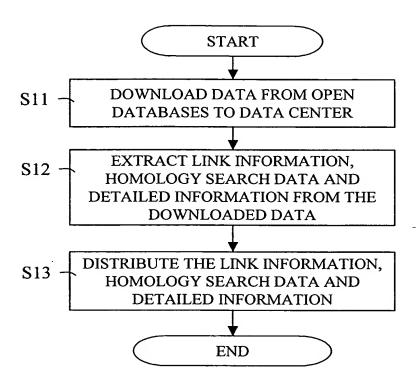
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FIG.1

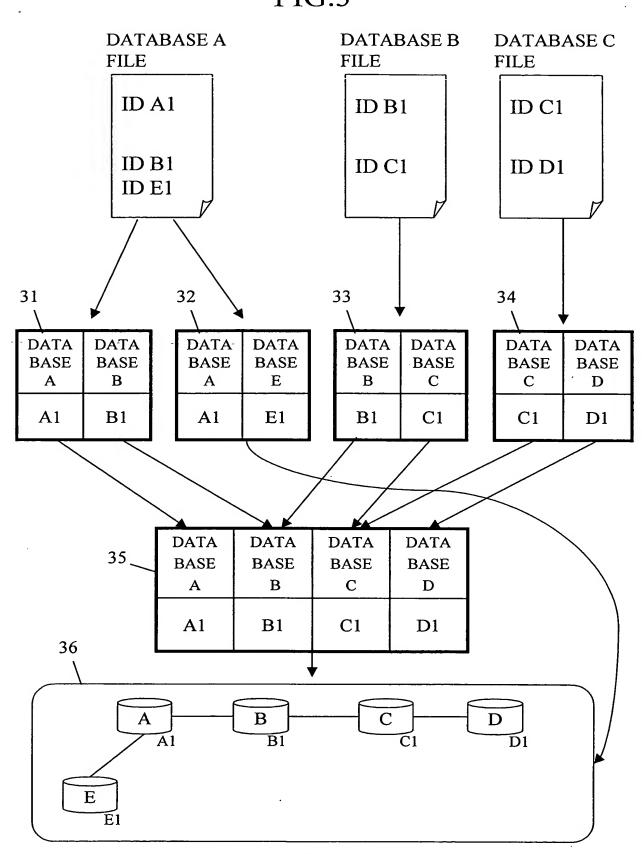


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FIG.2



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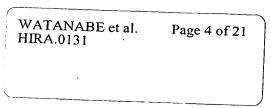
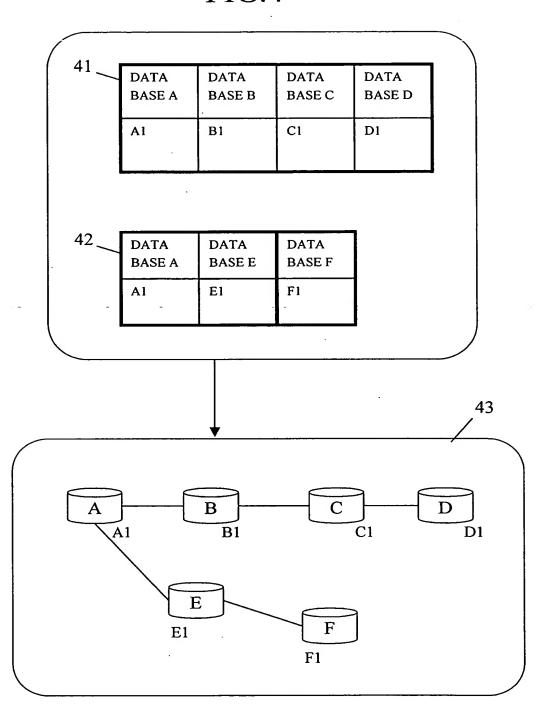


FIG.4



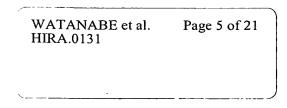
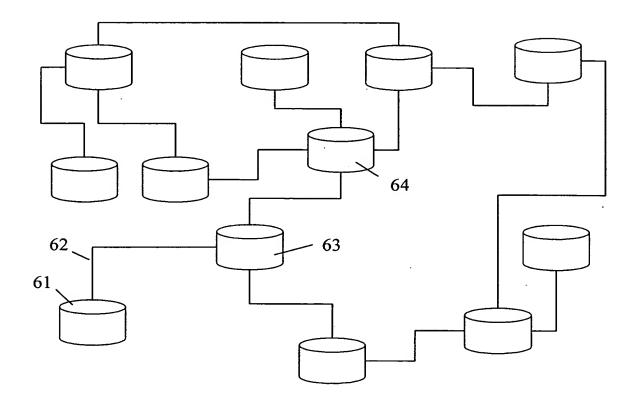


FIG.5

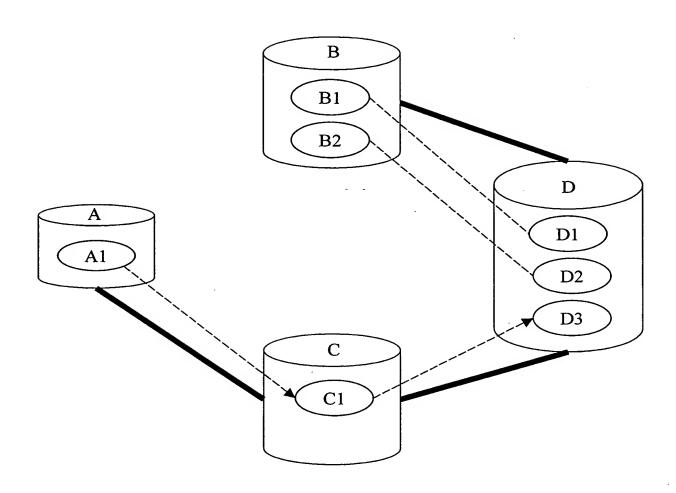
KeyDB	ROUTE	TargetDB
Α	В	С
Α	B, C	D
A	B, C	Е
	• • • •	

FIG.6



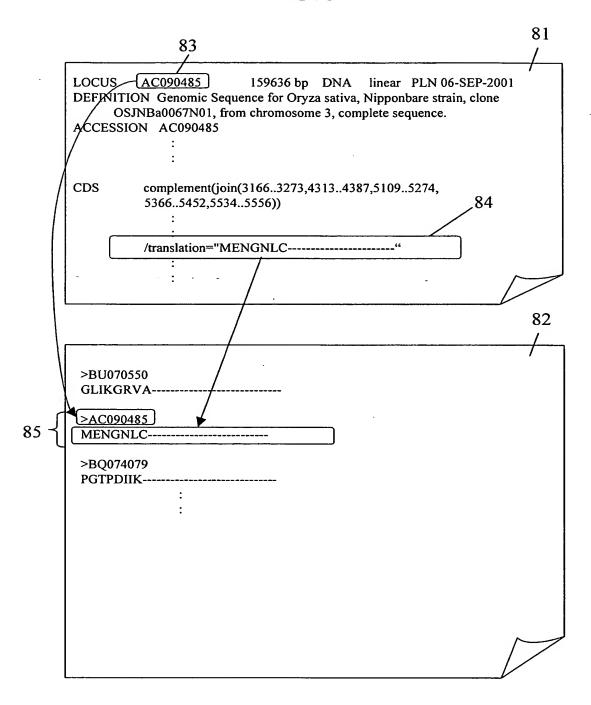
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FIG.7



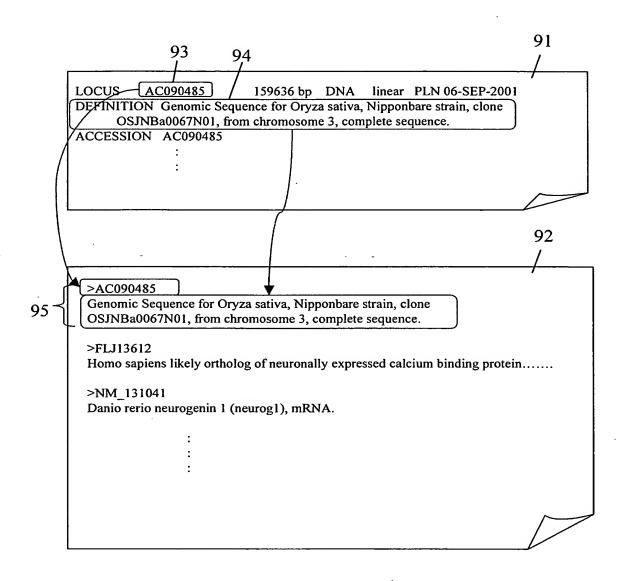
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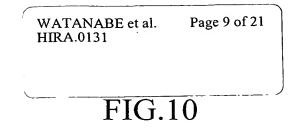
FIG.8

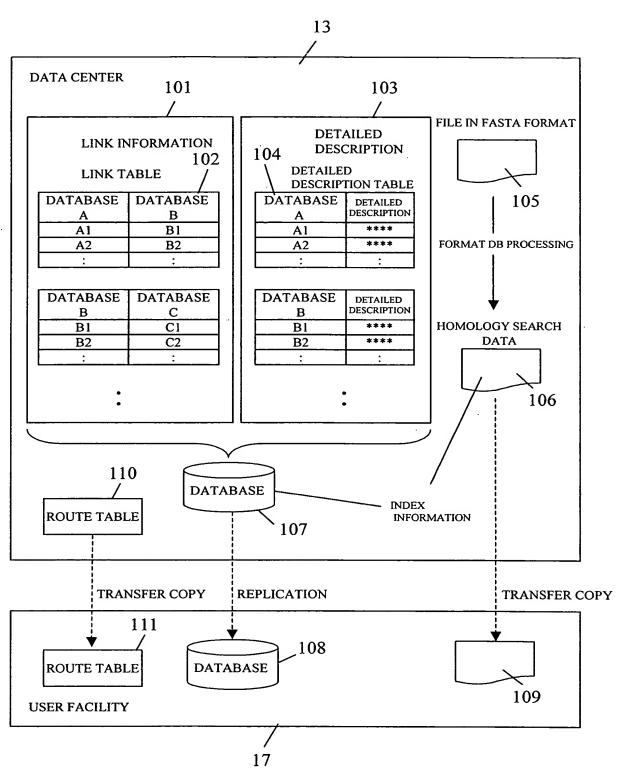


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FIG.9

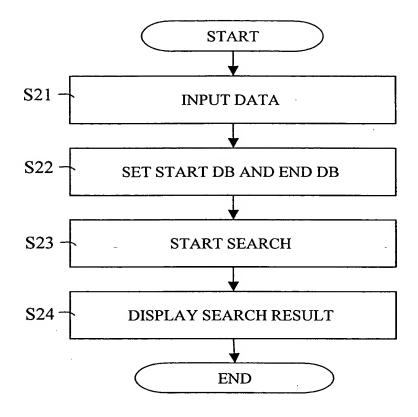






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FIG.11



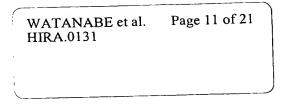
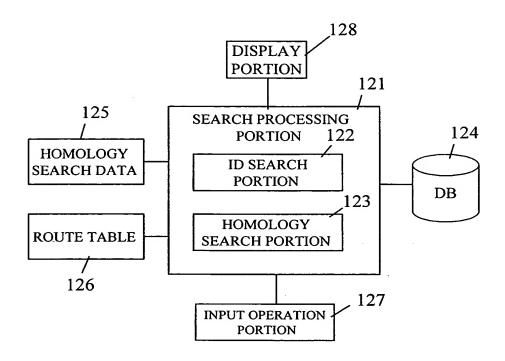
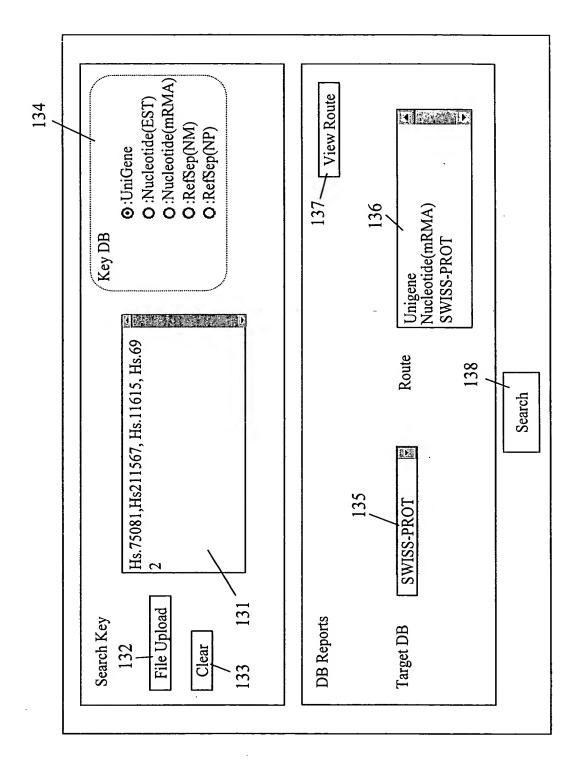
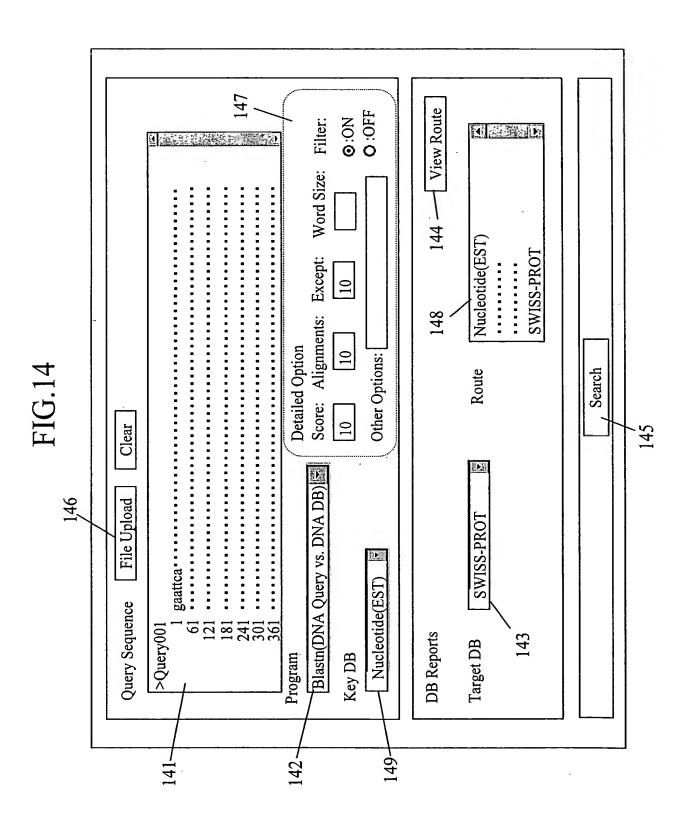


FIG.12









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FIG.15

Hs.0005 Hs.0012 Hs.0043 Hs.11706 Hs.75081

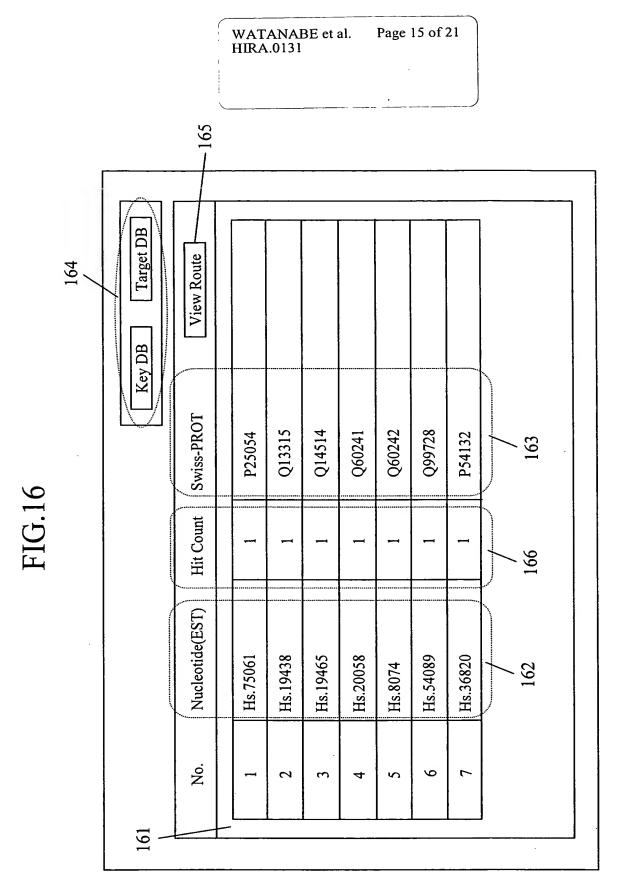


FIG.17

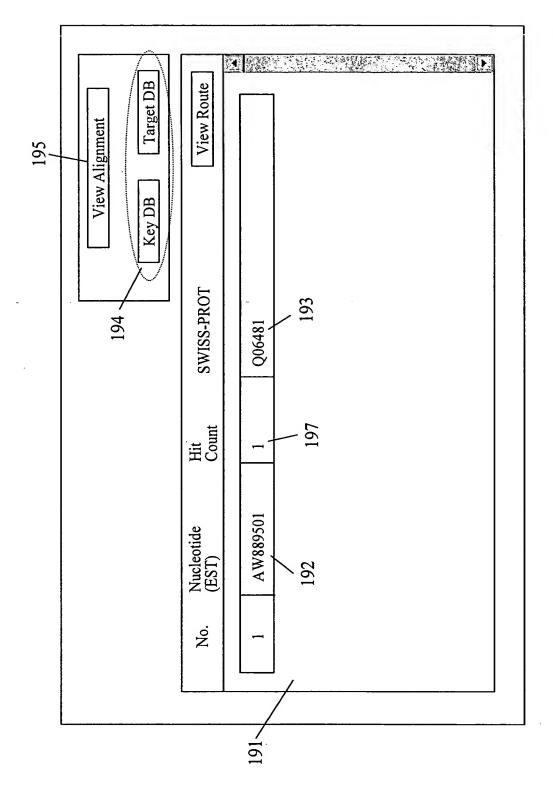
save close	Definition	601274939F1 NIH_MGC_20 Homo sapiens cDNA clone IMAGE:3615944 5', mRNA sep 602712606F1 NIH_MGC_48 Homo sapiens cDNA clone IMAGE:4853112 5', mRNA sep 601121460F1 NIH_MGC_20 Homo sapiens cDNA clone IMAGE:2988734 5', mRNA sep 603084468F1 NIH_MGC_120 Homo sapiens cDNA clone IMAGE:5223593 5', mRNA sep 60186366F1 NIH_MGC_17 Homo sapiens cDNA clone IMAGE:4845703 5', mRNA sep 602709227F1 NIH_MGC_19 Homo sapiens cDNA clone IMAGE:3613047 5', mRNA sep 601671261F1 NIH_MGC_10 Homo sapiens cDNA clone IMAGE:3183031 5', mRNA sep 601671261F1 NIH_MGC_20 Homo sapiens cDNA clone IMAGE:3954159 5', mRNA sep 602707881F1 NIH_MGC_43 Homo sapiens cDNA clone IMAGE:4844444 5', mRNA sep 603068266F1 NIH_MGC_118 Homo sapiens cDNA clone IMAGE:4844444 5', mRNA sep 603068266F1 NIH_MGC_118 Homo sapiens cDNA clone IMAGE:5216992 5', mRNA sep
Database: Nucleotide(FST	Accession	BE385140 BG758408 BE275447 BI834414 BF205012 BG750393 BE260149 BI758989 BF027028 BG749392 BI910571
Database: Ni	Search Key	Hs.350068 Hs.350068 Hs.350068 Hs.350068 Hs.350068 Hs.350068 Hs.350068 Hs.350068 Hs.350068

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FIG.18

> Query001			
1 gaattcca		 	
61		 	
121	<del></del>	 	
181		 	



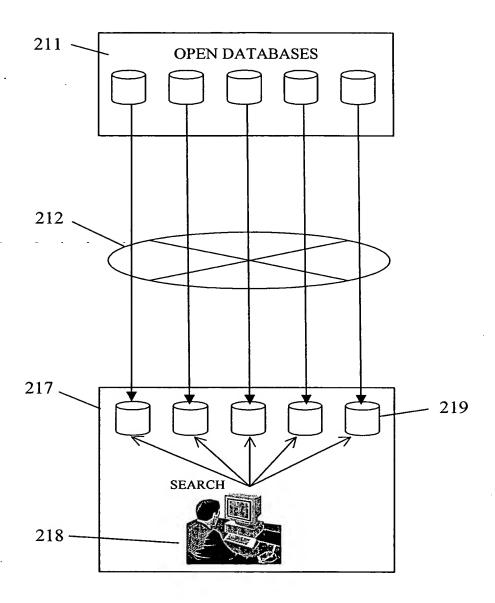


## FIG.20

Sequences producing significant alignments:	Score(bits	Score(bits) E value
EST   BF570078   602186096T1 NIH HGC 45 Homo sapiens cDNA clone	595	e-168
EST   BQ917841   AGENCOURT 8842358 Lupski sciatic nerve Hom	575	e-162
EST   BQ774659   UI-H-FHO-bcc-p-09-0-UI.s1 NCI_C GAP_FHO Ho	575	e-162
EST   BQ773379   UI-H-FEO-bby-c-13-0-UI.s1 NCI_C GAP_FEO Ho	575	e-162
EST   BQ773190   UI-H-FEO-bbv-j-22-0-UI.s2 NCI C GAP FEO Ho	575	e-162
EST   BQ772786   UI-H-FEO-bbu-i-13-0-UI.s2 NCI_C GAP_FEO Ho	575	e-162
EST   BQ772734   UI-H-FEO-bbr-m-04-0-UI.s1 NCI_C GAP_FEO Ho	575	e-162
EST   BQ668365   AGENCOURT 8303633 NIH HGC 102 Homo s	575	e-162
EST   BQ581190   il05h01.y1 Human insulinoma Homo sapiens cDNA	575	e-162
EST   BQ549102   ik88f09.y1 Human insulinoma Homo sapiens cDNA	575	e-162
>EST   BF570078   602186096T1 NIH_HGC_45 Homo sapiens cDNA clone IMAGE : 4310639 3' Length = 1040	GE: 431063	39 3' .
Score = $595 \text{ bits } (300)$ , Expect = $e-168$		
Identities = 300/300 (100%), Gaps = 1/298 (0%)		
Strand = Plus / Plus		
Query: 1 tgaagct / IllIII		
Sbict: 109		
•		
Sbjct: 169		

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FIG.21



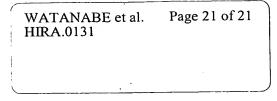


FIG.22

